0590
0823

#15

OIPE

RAW SEQUENCE LISTING DATE: 08/22/2002
PATENT APPLICATION: US/09/954,987B TIME: 15:57:20

Input Set : A:\seqlist.txt
Output Set: N:\CRF3\08222002\I954987B.raw

p.6

<110> APPLICANT: Stefan Bauer
Grayson B. Lipford
Hermann Wagner
<120> TITLE OF INVENTION: PROCESS FOR HIGH THROUGHPUT SCREENING OF
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<141> CURRENT FILING DATE: 2001-09-17
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PATENT APPLICATION: **US/09/954,987B** TIME: 15:57:21

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Output Set: **N:\CRF3\08222002\I954987B.raw**

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166 <213> ORGANISM: Mus musculus
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175  Phe Leu Lys Ser Val Pro Arg Phe Ser Ala Ala Ala Ser Cys Ser Asn
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182              100                 105                 110
183  Thr Ile Glu Pro Arg Thr Phe Leu Ala Met Arg Thr Leu Glu Glu Leu
184          115                 120                 125
185  Asn Leu Ser Tyr Asn Gly Ile Thr Thr Val Pro Arg Leu Pro Ser Ser
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187  Leu Val Asn Leu Ser Leu Ser His Thr Asn Ile Leu Val Leu Asp Ala
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189  Asn Ser Leu Ala Gly Leu Tyr Ser Leu Arg Val Leu Phe Met Asp Gly
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192              180                 185                 190
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195  Asn Asn Leu Thr Lys Val Pro Arg Gln Leu Pro Pro Ser Leu Glu Tyr
196      210                 215                 220
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202              260                 265                 270
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209  Tyr Glu Ser Ile Asn His Thr Asn Ala Phe Gln Asn Leu Thr Arg Leu
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217  Arg Trp Leu Ala Asp Leu Pro Lys Leu His Thr Leu His Leu Gln Met
218  385                 390                 395                 400
219  Asn Phe Ile Asn Gln Ala Gln Leu Ser Ile Phe Gly Thr Phe Arg Ala
220                  405                 410                 415
221  Leu Arg Phe Val Asp Leu Ser Asp Asn Arg Ile Ser Gly Pro Ser Thr
222              420                 425                 430
223  Leu Ser Glu Ala Thr Pro Glu Glu Ala Asp Asp Ala Glu Gln Glu Glu
224          435                 440                 445
225  Leu Leu Ser Ala Asp Pro His Pro Ala Pro Leu Ser Thr Pro Ala Ser
226      450                 455                 460
227  Lys Asn Phe Met Asp Arg Cys Lys Asn Phe Lys Phe Thr Met Asp Leu
228  465                 470                 475                 480
229  Ser Arg Asn Asn Leu Val Thr Ile Lys Pro Glu Met Phe Val Asn Leu
230                  485                 490                 495
231  Ser Arg Leu Gln Cys Leu Ser Leu Ser His Asn Ser Ile Ala Gln Ala
232              500                 505                 510
233  Val Asn Gly Ser Gln Phe Leu Pro Leu Thr Asn Leu Gln Val Leu Asp
234          515                 520                 525
235  Leu Ser His Asn Lys Leu Asp Leu Tyr His Trp Lys Ser Phe Ser Glu
236      530                 535                 540
237  Leu Pro Gln Leu Gln Ala Leu Asp Leu Ser Tyr Asn Ser Gln Pro Phe
238  545                 550                 555                 560
239  Ser Met Lys Gly Ile Gly His Asn Phe Ser Phe Val Ala His Leu Ser
240                  565                 570                 575
241  Met Leu His Ser Leu Ser Leu Ala His Asn Asp Ile His Thr Arg Val
242              580                 585                 590
243  Ser Ser His Leu Asn Ser Asn Ser Val Arg Phe Leu Asp Phe Ser Gly
244          595                 600                 605
245  Asn Gly Met Gly Arg Met Trp Asp Glu Gly Gly Leu Tyr Leu His Phe
246      610                 615                 620
247  Phe Gln Gly Leu Ser Gly Leu Leu Lys Leu Asp Leu Ser Gln Asn Asn
248  625                 630                 635                 640
249  Leu His Ile Leu Arg Pro Gln Asn Leu Asp Asn Leu Pro Lys Ser Leu
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251  Lys Leu Leu Ser Leu Arg Asp Asn Tyr Leu Ser Phe Phe Asn Trp Thr
252              660                 665                 670
253  Ser Leu Ser Phe Leu Pro Asn Leu Glu Val Leu Asp Leu Ala Gly Asn
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255  Gln Leu Lys Ala Leu Thr Asn Gly Thr Leu Pro Asn Gly Thr Leu Leu
256      690                 695                 700
257  Gln Lys Leu Asp Val Ser Ser Asn Ser Ile Val Ser Val Val Pro Ala
258  705                 710                 715                 720
259  Phe Phe Ala Leu Ala Val Glu Leu Lys Glu Val Asn Leu Ser His Asn
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Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

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Seq#:10; Xaa Pos. 39
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Seq#:145; Xaa Pos. 2,3,5,6,7,8,9,10,12,13
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